Ref #	Hits	Search Query	DBs	Default Operator	Plurals	Time Stamp
L1	8229	(data near5 mining)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2006/08/17 13:32
L2	4749	L1 and @ad<"20030101"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2006/08/17 13:35
L3	1324	L2 and "707".clas.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR .	OFF	2006/08/17 13:34
L4	611	L2 and ((regulation or decision) with tree\$1)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2006/08/17 13:35
L5	238	L4 and "707".clas.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2006/08/17 13:39
L6	201	L4 and 707/1-100.ccls.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2006/08/17 13:48
L7	39	L5 and 707/5.ccls.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2006/08/17 13:38
L8	44	L5 and 707/100.ccls.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2006/08/17 13:38

L9	123	L4 and "706".clas.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2006/08/17 14:03
L10	26	L4 and 707/1.ccls.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2006/08/17 13:41
L11	56	L4 and 707/6.ccls.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2006/08/17 13:42
L12	0	L4 and 707/47.ccls.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2006/08/17 13:43
L13	0	L1 and 707/47.ccls.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2006/08/17 13:43
L14	1	L1 and 370/255.ccls.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2006/08/17 13:43
L15	2	"20040267770"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2006/08/17 13:48
L16	44	L4 and 707/100.ccls.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2006/08/17 13:49

L17	53	L4 and 707/102.ccls.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2006/08/17 13:50
L18	3	L4 and 707/200.ccls.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2006/08/17 13:50
L19	92	L4 and "705".clas.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2006/08/17 14:09
L20	1	L4 and 705/35.ccls.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2006/08/17 14:05
L21	0	L4 and 370/255.ccls.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2006/08/17 14:05
L22	1	L2 and 370/255.ccls.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2006/08/17 14:05
L23	439	(data near5 mining) with (visual or visualization\$1)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2006/08/17 14:09
S1	1580082	computer	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 11:08

S2	6	"609490".ap.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 14:07
S3	5	"423011".ap.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/28 13:26
S4	5064	(decision adj tree\$1)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 16:28
S5	725	S4 and (data adj mining)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/28 15:48
S6	407	S5 and (interactive or dynamic)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/28 15:12
S7		S6 and (configuration adj tree\$1)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/28 15:00
S8	403	configuration adj tree\$1	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/28 15:01
S9	5	S8 and (data adj mining)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/28 15:01

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S10	3	"423678".ap.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 14:34
S11	2	"423678".ap. and ((data adj mining) or tree or decision or configuration or model or modeling or interactive or user or dynamic or integration or rank or ranking or update or learn or learning or analyze or analysis or visualize or visualization)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/28 15:24
S12	1	"972057".ap.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/28 15:24
S13	16	regulation adj tree\$1	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/28 15:34
S14	6	"609490".ap.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 11:35
S15	16	regulation adj tree\$1	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/28 15:37
S16	30704	S15 and "706"/.clas. or "707"/.clas.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/28 15:37
S17	3	S15 and "706"/.clas.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/28 15:37

S18	1	S15 and "707"/.clas.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/28 15:37
S19	564	S5 and @ad<"20030625"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/28 15:48
S20	170	S19 and (rank or ranking)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR .	OFF	2005/11/28 15:48
S21	3	S20 and (rule adj ranking)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR ·	OFF	2005/11/28 16:08
S22	5064	decision adj tree\$1	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF .	2005/11/28 16:08
S23	156	S22 and lee.in.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/28 16:08
S24	1	S23 and (configuration adj tree)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	ÖR	OFF	2005/11/28 16:08
S25	0	5745601l.pn.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/29 11:19

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S26	2	"5745601".pn.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/29 11:21
S27·	16	regulation adj tree\$1	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/29 11:24
S28	1	"972,057".ap.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/29 11:48
S29	6	"081441".ap.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/29 11:48
S30	7	"118553".ap.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/29 13:45
S31	7	"104647".ap.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/29 13:46
S32	2	"104647".ap. and (model or modeling or modelling or tree or decision or regulation or rank or ranking or rule\$1 or mine or mining)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/29 13:51
S33		"972057".ap.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/29 15:01

						
S34	0	"5892801".pn. and berry	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/29 13:59
S35	2	"5892801".pn.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/29 13:59
S36	2	"6941304".pn.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/29 15:01
S37	2	"6941303".pn.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/29 15:05
S38	30	rule adj ranking	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/29 15:07
S39	. 10	S38 and (local and global)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/29 15:07
S40	7	S39 and (population)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/29 15:08
S41	0	S40 and statics	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/29 15:07

S42	2	S40 and statistics	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/29 15:07
S43	0	S42 and @ad<"20030625"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/29 15:08
S44	5	S40 and @ad<"20030625"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR .	OFF	2005/11/29 15:08
S45	6063	data adj mining	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF [*]	2005/11/30 11:08
S46	4514	S45 and @ad<"20030625"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 11:09
S47	150	S46 and zoom	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 11:09
S48	9	S47 and (feature with zoom)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 11:09
S49	5	"747515".ap.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 11:44

S50	5069	Decision adj tree\$1	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR .	OFF	2005/11/30 11:45
S51	134625	"L9" and @ad<"20030625"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 11:45
S52	3982	S50 and @ad<"20030625"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 11:45
S53	212	S52 and (multilevel or multi-level)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 11:46
S54	31	S53 and ((decision adj tree\$1) with level\$1)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 13:20
S55	5	"957637".ap.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 12:06
S56	5069	Decision adj tree\$1	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 13:20
S57	3982	S56 and @ad<"20030625"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 13:20

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S58	212	S57 and (multilevel or multi-level)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 13:20
S59	31	S58 and ((decision adj tree\$1) with level\$1)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 13:20
S60	0	S59 and (contrast with example)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 13:22
S61	1385	(model near3 (present or presenting or presentation))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 13:23
S62	1084	S61 and @ad<"20030625"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 13:23
S63	334	S62 and contrast	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 13:23
S64	338	S62 and (contrast or contrasting)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 13:23
S65	290	S64 and (group\$1 or grouping)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 13:24

S66	41	S65 and (rank or ranks or ranking)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 13:30
S67	0	S66 and ((contrast\$4 with example\$1) and (rank\$4 same group\$5) and label\$1)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 13:26
S68	0	S66 and (contrast\$4 with example\$1) and (rank\$4 same group\$5)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 13:26
S69	0	contrast adj example\$1	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 13:32
S70	5069	decision adj tree\$1	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 13:32
S71	3982	S70 and @ad<"20030625"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 13:33
S72	23	S71 and (present\$5 with contrast\$4)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 13:33
S73	7	"854301".ap.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 14:10

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S74	2	"6826552".pn.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 14:17
S75	2	"5878406".pn.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 14:18
S76	3	(similar adj (characteristics or feature\$1 or qualities)) with (different near3 label\$1)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 14:20
S77	0	S76 and @ad<"20030625"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 14:20
S78	3	(similar near3 (characteristics or feature\$1 or qualities)) with (different near3 label\$1)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR .	OFF	2005/11/30 14:21
S79	73	contrast\$4 with example\$1	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 14:21
S80	7	S79 and @ad<"20030625"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 14:23
581	226084	similar with different	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 14:23

S82	140177	S81 and @ad<"20030625"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 14:23
S83	1	S82 and (differnt adj label\$1)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 14:24
S84	34594	contrasting	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 14:25
S85	21409	S84 and @ad<"20030625"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 14:25
S86	44	S85 and (contrasting with (similar with different))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 14:30
S87	0	6941303l.pn.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 14:30
S88	2	"6941303".pn.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 14:30
S89	93611	(output with group\$5)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 14:35

S90	73669	S89 and @ad<"20030625"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 14:35
S91	2326	S90 and (group\$5 with present\$5)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 14:36
S92	668	S91 and (model\$1 or modeling)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 16:04
S93	121302	normalize\$1	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 16:05
S94	98677	S93 and @ad<"20030625"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 16:05
S95 _.	1895	S94 and "707"/.clas.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 16:27
S96	2	"6941303".pn.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR .	OFF	2005/11/30 16:27
S97	5069	(decision adj tree\$1)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 16:28

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S98	3982	S97 and @ad<"20030625"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR ·	OFF	2005/11/30 18:18
S99	13363	(remove\$1 with node\$1)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 16:29
S10 0	89	S98 and (remove\$1 with node\$1)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 16:29
S10 1	83	S100 and add\$3	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 16:30
S10 2	65	S101 and updat\$4	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 16:30
S10 3	3	S102 and (remov\$4 near3 rule)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 16:48
S10 4	1	"972057".ap.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 16:36
S10 5	2	S102 and (remov\$4 near3 sample\$1)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 16:49

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S10 6	5069	(decision adj tree\$1)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 18:18
S10 7	3982	S106 and @ad<"20030625"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 18:18
\$10 8	288	S107 and histogram	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 18:19
S10 9	0	S108 and (histograqm with bar\$1)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 18:20
S11 0	21	S108 and (histogram with bar\$1)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 18:44
S11 1	7	"813,336".ap.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 18:41
S11 2	7	"815,473".ap.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 18:41
S11 3		S108 and (histogram same bar\$1)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 19:04

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S11 6	273	highlight\$1 near represent\$5	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 19:07
S11 7	210	S116 and @ad<"20030625"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 19:07
S11 8	6	S117 and "707"/.clas.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	OFF	2005/11/30 19:22
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1 Bioinformatics (BIO): Incremental interactive mining of constrained association rules



from biological annotation data with nominal features

Imad Rahal, Dongmei Ren, Amal Perera, Hassan Najadat, William Perrizo, Riad Rahhal, Willy Valdivia

March 2005 Proceedings of the 2005 ACM symposium on Applied computing SAC '05

Publisher: ACM Press

Full text available: 完 pdf(132.24 KB) Additional Information: full citation, abstract, references, index terms

Data arising from genomic and proteomic experiments is amassing at high speeds resulting in huge amounts of raw data; consequently, the need for analyzing such biological data --- the understanding of which is still lagging way behind --- has been prominently solicited in the post-genomic era we are currently witnessing. In this paper we attempt to analyze annotated genome data by applying a very central data-mining technique known as association rule mining with the aim of discovering rules cap ...

Keywords: P-trees, association rule mining, bioinformatics, incremental, interactive, yeast genome

2 Contributed articles on online, interactive, and anytime data mining: MobiMine:





monitoring the stock market from a PDA

Hillol Kargupta, Byung-Hoon Park, Sweta Pittie, Lei Liu, Deepali Kushraj, Kakali Sarkar January 2002 ACM SIGKDD Explorations Newsletter, Volume 3 Issue 2

Publisher: ACM Press

Full text available: 包 pdf(1.16 MB)

Additional Information: full citation, abstract, references, citings

This paper describes an experimental mobile data mining system that allows intelligent monitoring of time-critical financial data from a hand-held PDA. It presents the overall system architecture and the philosophy behind the design. It explores one particular aspect of the system---automated construction of personalized focus area that calls for user's attention. This module works using data mining techniques. The paper describes the data mining component of the system that employs a novel Four ...

3 Link mining applications: progress and challenges



Ted E. Senator

December 2005 ACM SIGKDD Explorations Newsletter, Volume 7 Issue 2

Publisher: ACM Press

Full text available: Additional Information: full citation, abstract, references, index terms

This article reviews a decade of progress in the area of link mining, focusing on application requirements and how they have and have not yet been addressed, especially in the area of complex event detection. It discusses some ongoing challenges and

suggests ideas that could be opportunities for solutions. The most important conclusion of this article is that while there are many link mining techniques that work well for individual link mining tasks, there is not yet a comprehensive framework th ...

Keywords: complex event detection, data mining applications, link analysis, link discovery, link mining, pattern analysis, pattern discovery, pattern matching, structured data

4 Articles on microarray data mining: Towards interactive exploration of gene



expression patterns

Daxin Jiang, Jian Pei, Aidong Zhang

December 2003 ACM SIGKDD Explorations Newsletter, Volume 5 Issue 2

Publisher: ACM Press

Full text available: pdf(527.68 KB) Additional Information: full citation, abstract, references

Analyzing coherent gene expression patterns is an important task in bioinformatics research and biomedical applications. Recently, various clustering methods have been adapted or proposed to identify clusters of co-expressed genes and recognize coherent expression patterns as the centroids of the clusters. However, the interpretation of coexpressed genes and coherent patterns mainly depends on the domain knowledge, which presents several challenges for coherent pattern mining and cannot be solv ...

5 E-government services and policy track: An e-government information architecture



for regulation analysis and compliance assistance

Gloria T. Lau, Shawn Kerrigan, Kincho H. Law, Gio Wiederhold March 2004 Proceedings of the 6th international conference on Electronic commerce **ICEC '04**

Publisher: ACM Press

Full text available: pdf(510.70 KB)

Additional Information: full citation, abstract, references, citings, index terms

The complexity and diversity of government regulations make understanding the regulations a non-trivial task. One of the issues is the existence of multiple sources of regulations and interpretive guides. In this work, we propose an information infrastructure for regulation analysis, which includes a document repository and tools for compliance assistance and similarity analysis. A regulatory repository is developed based on an XML format, and important features, such as concepts and measurement ...

Keywords: compliance check, e-government, e-rulemaking, legal informatics, shallow parsing, similarity analysis, text mining

6 KDD-99 conference reports: Knowledge discovery in databases: 10 years after





Gregory Piatetsky-Shapiro January 2000 ACM SIGKDD Explorations Newsletter, Volume 1 Issue 2

Publisher: ACM Press

Full text available: 党 pdf(264.76 KB) Additional Information: full citation, abstract, references

In this paper, we describe the past IO years of KDD and outline predictions for the next 10 years.

Keywords: KDD, data mining, history, knowledge discovery in databases

7 Industry/government track paper: Using relational knowledge discovery to prevent securities fraud



Jennifer Neville, Özgür Şimşek, David Jensen, John Komoroske, Kelly Palmer, Henry Goldberg

August 2005 Proceeding of the eleventh ACM SIGKDD international conference on

Knowledge discovery in data mining KDD '05

Publisher: ACM Press

Full text available: <u>同 pdf(1.22 MB)</u> Additional Information: <u>full citation</u>, abstract, references, index terms

We describe an application of relational knowledge discovery to a key regulatory mission of the National Association of Securities Dealers (NASD). NASD is the world's largest private-sector securities regulator, with responsibility for preventing and discovering misconduct among securities brokers. Our goal was to help focus NASD's limited regulatory resources on the brokers who are most likely to engage in securities violations. Using statistical relational learning algorithms, we developed mod ...

Keywords: fraud detection, relational probability trees, statistical relational learning

8 Industry track papers: Handling very large numbers of association rules in the

analysis of microarray data

Alexander Tuzhilin, Gediminas Adomavicius

July 2002 Proceedings of the eighth ACM SIGKDD international conference on Knowledge discovery and data mining

Publisher: ACM Press

Full text available: pdf(953.83 KB)

Additional Information: full citation, abstract, references, citings, index terms

The problem of analyzing microarray data became one of important topics in bioinformatics over the past several years, and different data mining techniques have been proposed for the analysis of such data. In this paper, we propose to use association rule discovery methods for determining associations among expression levels of different genes. One of the main problems related to the discovery of these associations is the scalability issue. Microarrays usually contain very large numbers of genes ...

Keywords: analysis of microarray data, association rules, bioinformatics, expert-driven rule validation, post-processing of discovered rules, rule filtering, rule grouping

9 Industrial/government track: Similarity analysis on government regulations

Gloria T. Lau, Kincho H. Law, Gio Wiederhold

August 2003 Proceedings of the ninth ACM SIGKDD international conference on Knowledge discovery and data mining

Publisher: ACM Press

Full text available: Topdf(254.75 KB) Additional Information: full citation, abstract, references, index terms

Government regulations are semi-structured text documents that are often voluminous, heavily cross-referenced between provisions and even ambiguous. Multiple sources of regulations lead to difficulties in both understanding and complying with all applicable codes. In this work, we propose a framework for regulation management and similarity analysis. An online repository for legal documents is created with the help of text mining tool, and users can access regulatory documents either through the ...

Keywords: legal informatics, regulations, similarity analysis, text mining

10 Assessment by belief

Takao Miura, Isamu Shioya

January 2001 Proceedings of the 12th Australasian database conference ADC '01

Publisher Site

Additional Information: full citation, abstract, references, index terms

We discuss how to post-evaluate inductive classification based on users belief. Although we could learn classification rules inductively by means of decision tree generation, we wonder whether it is consistent with our utilization or not. In this investigation we discuss



how to obtain assessment of learning results by verifying belief. Our idea is based on Decision Tree with Hierarchy to class and attributes; to each attribute we assume taxonomy on the domain in addition to class hierarchy. Then ...

Keywords: knowledge discovery, machine Learning, mining in databases

11 Nonorthogonal decomposition of binary matrices for bounded-error data compression



and analysis

Mehmet Koyutürk, Ananth Grama, Naren Ramakrishnan

March 2006 ACM Transactions on Mathematical Software (TOMS), Volume 32 Issue 1

Publisher: ACM Press

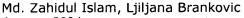
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Additional Information: full citation, abstract, references, index terms

This article presents the design and implementation of a software tool, PROXIMUS, for error-bounded approximation of high-dimensional binary attributed datasets based on nonorthogonal decomposition of binary matrices. This tool can be used for analyzing data arising in a variety of domains ranging from commercial to scientific applications. Using a combination of innovative algorithms, novel data structures, and efficient implementation, PROXIMUS demonstrates excellent accuracy, performance, and ...

Keywords: Compressing binary-valued vectors, nonorthogonal matrix decompositions, semidiscrete decomposition

12 A framework for privacy preserving classification in data mining



January 2004 Proceedings of the second workshop on Australasian information security, Data Mining and Web Intelligence, and Software **Internationalisation - Volume 32 ACSW Frontiers '04**

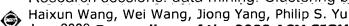
Publisher: Australian Computer Society, Inc.

Full text available: 🔁 pdf(365.56 KB) Additional Information: full citation, abstract, references

Nowadays organizations all over the world are dependent on mining gigantic datasets. These datasets typically contain delicate individual information, which inevitably gets exposed to different parties. Consequently privacy issues are constantly under the limelight and the public dissatisfaction may well threaten the exercise of data mining and all its benefits. It is thus of great importance to develop adequate security techniques for protecting confidentiality of individual values used for dat ...

Keywords: data mining, data security, noise addition, privacy, statistical database

13 Research sessions: data mining: Clustering by pattern similarity in large data sets



June 2002 Proceedings of the 2002 ACM SIGMOD international conference on Management of data SIGMOD '02

Publisher: ACM Press

Full text available: 完 pdf(1.09 MB)

Additional Information: full citation, abstract, references, citings, index terms

Clustering is the process of grouping a set of objects into classes of *similar* objects. Although definitions of similarity vary from one clustering model to another, in most of these models the concept of similarity is based on distances, e.g., Euclidean distance or cosine distance. In other words, similar objects are required to have close values on at least a set of dimensions. In this paper, we explore a more general type of similarity. Under the pCluster model we proposed, two objects ...

14 Biclustering Algorithms for Biological Data Analysis: A Survey Sara C. Madeira, Arlindo L. Oliveira



January 2004 IEEE/ACM Transactions on Computational Biology and Bioinformatics (TCBB), Volume 1 Issue 1

Publisher: IEEE Computer Society Press

Additional Information: full citation, references, citings

Keywords: Biclustering, simultaneous clustering, coclustering, subspace clustering, bidimensional clustering, direct clustering, block clustering, two-way clustering, two-mode clustering, two-sided clustering, microarray data analysis, biological data analysis, gene expression data.

15 Data mining (DM): A hybrid approach for multiresolution modeling of large-scale



scientific data

Tina Eliassi-Rad, Terence Critchlow

March 2005 Proceedings of the 2005 ACM symposium on Applied computing SAC '05

Publisher: ACM Press

Full text available: pdf(428.94 KB) Additional Information: full citation, abstract, references, index terms

Simulations of complex scientific phenomena involve the execution of massively parallel computer programs. These simulation programs generate large-scale multidimensional data sets over the spatio-temporal region. Analyzing such massive data sets is an essential step in helping scientists glean new information. To this end, efficient and effective data models are needed. In this paper, we present a hybrid approach for constructing data models from large-scale multidimensional scientific data set ...

Keywords: information retrieval, large-scale scientific data sets, multiresolution indices, multivariate clusters, topological models

16 Research track papers: Turning CARTwheels: an alternating algorithm for mining redescriptions





Naren Ramakrishnan, Deept Kumar, Bud Mishra, Malcolm Potts, Richard F. Helm August 2004 Proceedings of the tenth ACM SIGKDD international conference on Knowledge discovery and data mining KDD '04

Publisher: ACM Press

Additional Information: full citation, abstract, references, citings, index terms

We present an unusual algorithm involving classification trees---CARTwheels---where two trees are grown in opposite directions so that they are joined at their leaves. This approach finds application in a new data mining task we formulate, called redescription mining. A redescription is a shift-of-vocabulary, or a different way of communicating information about a given subset of data; the goal of redescription mining is to find subsets of data that afford multiple descriptions. We highli ...

Keywords: classification trees, data mining in biological domains, redescriptions

17 Multi Relational Data Mining (MRDM): Biological applications of multi-relational data



mining

David Page, Mark Craven

July 2003 ACM SIGKDD Explorations Newsletter, Volume 5 Issue 1

Publisher: ACM Press

Additional Information: full citation, abstract, references, citings

Biological databases contain a wide variety of data types, often with rich relational structure. Consequently multi-relational data mining techniques frequently are applied to biological data. This paper presents several applications of multi-relational data mining to biological data, taking care to cover a broad range of multi-relational data mining

techniques.

18 Research papers: mining biological and medical data: TRICLUSTER: an effective algorithm for mining coherent clusters in 3D microarray data



Lizhuang Zhao, Mohammed J. Zaki

June 2005 Proceedings of the 2005 ACM SIGMOD international conference on Management of data

Publisher: ACM Press

Full text available: 🛱 pdf(466.52 KB) Additional Information: full citation, abstract, references

In this paper we introduce a novel algorithm called TRICLUSTER, for mining coherent clusters in three-dimensional (3D) gene expression datasets. TRICLUSTER can mine arbitrarily positioned and overlapping clusters, and depending on different parameter values, it can mine different types of clusters, including those with constant or similar values along each dimension, as well as scaling and shifting expression patterns. TRICLUSTER relies on graph-based approach to mine all valid clusters. For eac ...

19 Data mining, knowledge discovery & OLTP: Using secure coprocessors for privacy



preserving collaborative data mining and analysis

Bishwaranjan Bhattacharjee, Naoki Abe, Kenneth Goldman, Bianca Zadrozny, Vamsavardhana R. Chillakuru, Marysabel del Carpio, Chid Apte

June 2006 Proceedings of the 2nd international workshop on Data management on new hardware DaMoN '06

Publisher: ACM Press

Full text available: pdf(248.64 KB) Additional Information: full citation, abstract, references, index terms

Secure coprocessors have traditionally been used as a keystone of a security subsystem, eliminating the need to protect the rest of the subsystem with physical security measures. With technological advances and hardware miniaturization they have become increasingly powerful. This opens up the possibility of using them for non traditional use. This paper describes a solution for privacy preserving data sharing and mining using cryptographically secure but resource limited coprocessors. It uses me ...

Keywords: collaboration, data mining, federation, privacy

20 Bioinformatics—an introduction for computer scientists



Jacques Cohen

June 2004 ACM Computing Surveys (CSUR), Volume 36 Issue 2

Publisher: ACM Press

Full text available: Ref pdf(261.56 KB) Additional Information: full citation, abstract, references, index terms

The article aims to introduce computer scientists to the new field of bioinformatics. This area has arisen from the needs of biologists to utilize and help interpret the vast amounts of data that are constantly being gathered in genomic research---and its more recent counterparts, proteomics and functional genomics. The ultimate goal of bioinformatics is to develop in silico models that will complement in vitro and in vivo biological experiments. The article provides a bird's eye view of the ...

Keywords: DNA, Molecular cell biology, RNA and protein structure, alignments, cell simulation and modeling, computer, dynamic programming, hidden-Markov-models, microarray, parsing biological sequences, phylogenetic trees

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